



## A super-resolution approach for receptive fields estimation of neuronal ensembles

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### ► To cite this version:

Daniela Pamplona, Gerrit Hilgen, Sahar Pirmoradian, Matthias H. Hennig, Bruno Cessac, et al.. A super-resolution approach for receptive fields estimation of neuronal ensembles. Computational Neuroscience (CNS), Jul 2015, Prague, Czech Republic. . hal-01215541

**HAL Id: hal-01215541**

**<https://inria.hal.science/hal-01215541>**

Submitted on 19 Oct 2015

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# A super-resolution approach for receptive fields estimation of neuronal ensembles

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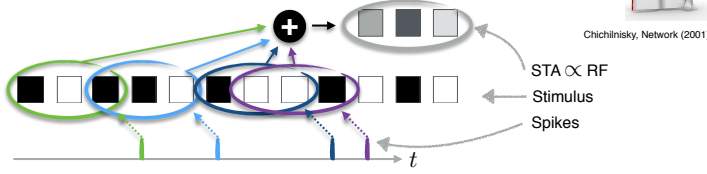
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## Spike Triggered Average (STA) with white noise (WN)

### A classical technique to find receptive fields (RF)



**Convergence:** Error tends to zero as the spike count tends to infinity.

$$\mathcal{E}(STA, RF) = \sum_{x,y,t} (STA(x,y,t) - RF(x,y,t))^2 \approx \mathcal{N}(0,1)^2 \frac{\sigma^2}{n(T)}$$

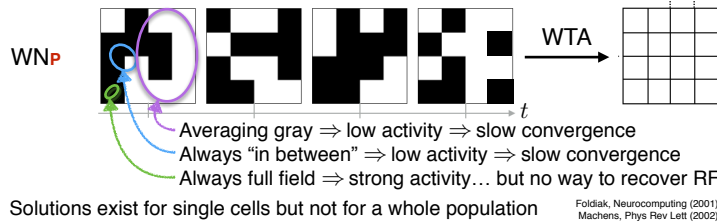
Gaussian distribution

Variance of the stimulus

Spike count

### Spatial resolution of the estimation

It is defined by the size of the blocks of the checkerboard stimulus.  
How to set block size? A matter of compromise...



## How to increase the resolution of RF over the population?

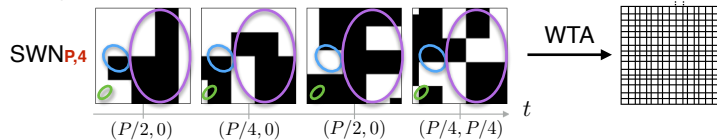
### Naïve solution: decrease block size



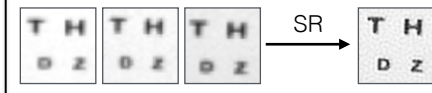
### Proposed solution: Introducing shifted white noise (SWN)

We define SWN as a WN stimulus where blocks are randomly moved by discrete spatial shifts at each time stamp.

Example with 4 shifts:



**Remark** Analogy with super-resolution (SR) techniques in image processing



Using SWN<sub>P,S</sub> increases the STA resolution by a factor **S** without decreasing the block size, thus it is suitable for a large ensemble of neurons.

The STA obtained can still be theoretically related to RF in the context of LN models similarly to the classical white noise scenario.

## Simulation study

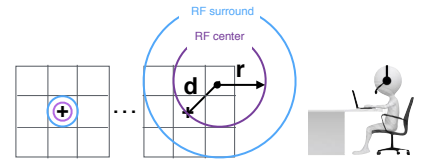
### Population of LNP neurons

Spatial RFs: DOG

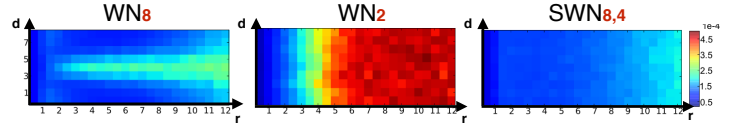
Temporal RFs: DOE

5 trials, 20000 images per trial

Display rate 30Hz



### STA error as a function of the cell RF position and size:



### Results

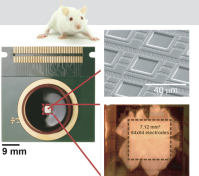
For a fixed block size, the STA spatial resolution is improved as a function of the number of possible shifts, leading to smaller estimation error.  
Targeting a given STA spatial resolution, shifting the stimulus increases the convergence speed.

(see COSYNE 2015 poster for more results)

## Experimental study

### Experimental protocol

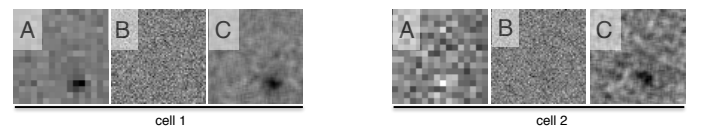
Pan-retinal RGC responses from WT mice were recorded using the high-density large-scale CMOS-based APS MEA (Biocam, 3Brain GmbH, CH) featuring 4096 electrodes (42  $\mu$ m spacing) arranged in a 64x64 configuration, covering an active area of 7.12 mm<sup>2</sup>.



White Noise images (block size of 160 $\mu$ m, 40 $\mu$ m and 160 $\mu$ m with arbitrary shifts of 40 $\mu$ m) were presented for 15 min at 30Hz (P79) and 15 Hz (P76) using a custom built system based on a DLP video projector.

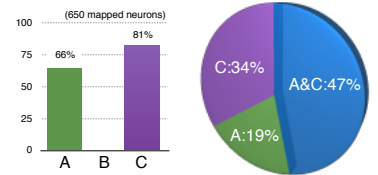
A: WN<sub>160</sub>  
B: WN<sub>40</sub>  
C: SWN<sub>160,40</sub>

### Comparison between RF estimations

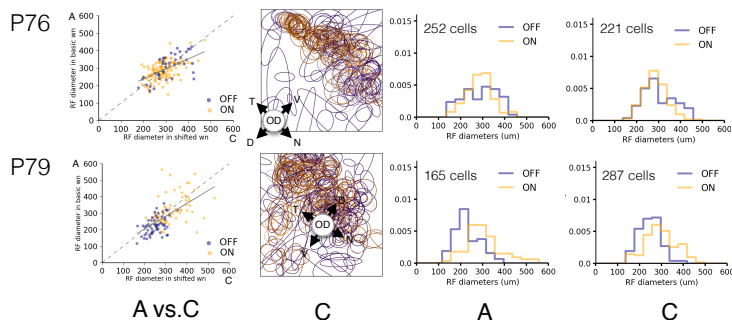


### Statistical analysis over the population

Most of RF are mapped by both stimuli (with highest resolution from SWN).  
Differences are possibly coming from RF properties (e.g., size and position)



### Distribution of RF sizes



WN and SWN give similar RF estimation. SWN allow to map more large RF.

This study presents for the first time a simple approach allowing quick and simultaneous extraction of receptive field profiles with great accuracy from hundreds to thousands of retinal ganglion cells. The method not only ensures strong responses because these stimuli are big (e.g. 160  $\mu$ m), but it also provides receptive field measurements at super resolution thanks to the small shifts (40  $\mu$ m) applied to the basic checkerboard stimuli.

This work was partially supported by the EC IP project FP7-ICT-2011-9 no. 600847 (RENVISION)

